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OIPE

RAW SEQUENCE LISTING DATE: 09/01/2004
PATENT APPLICATION: US/09/825,423C TIME: 17:20:04

Input Set : A:\ID01152 Sequence Listing.txt
Output Set: N:\CRF4\09012004\I825423C.raw

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3 <110> APPLICANT: Weber, Patricia C.
        Reichert, Paul
 5 .
        Madison, Vincent S.
 6
        Wyss, Daniel
 7
        Yao, Nanhua
 8
        Liu, Dingjiang
 9
        Gesell, Jennifer
11 <120> TITLE OF INVENTION: Hepatitis C Virus NS3 Helicase Fragments
13 <130> FILE REFERENCE: ID01152 US
15 <140> CURRENT APPLICATION NUMBER: US 09/825,423C
16 <141> CURRENT FILING DATE: 2001-04-03
18 <150> PRIOR APPLICATION NUMBER: US 60/194,419
19 <151> PRIOR FILING DATE: 2000-04-04
21 <160> NUMBER OF SEQ ID NOS: 16
23 <170> SOFTWARE: PatentIn Ver. 2.1
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 631
27 <212> TYPE: PRT
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34 Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu
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37 Val Gln Ile Val Ser Thr Ala Thr Gln Thr Phe Leu Ala Thr Cys Ile
40 Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile
43 Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp Gln
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                                            75
46 Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro
49 Cys Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp
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                                   105
52 Val Ile Pro Val Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser
           115
                               120
55 Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu
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58 Cys Pro Ala Gly His Ala Val Gly Leu Phe Arg Ala Ala Val Cys Thr
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61 Arg Gly Val Thr Lys Ala Val Asp Phe Ile Pro Val Glu Asn Leu Glu
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64 Thr Thr Met Arg Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala
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67 ۲	Val	Pro	Gln	Ser	Phe	Gln	Val	Ala	His	Leu	His	Ala	Pro	Thr	Gly	Ser
68			195					200					205		-	
70 0	Gly	Lys	Ser	Thr	Lys	Val	Pro	Ala	Ala	Tyr	Ala	Ala	Gln	Gly	Tyr	Lys
71	_	210			-		215			-		220		•	•	•
73 \	Val	Leu	Val	Leu	Asn	Pro	Ser	Val	Ala	Ala	Thr		Gly	Phe	Gly	Ala
74 2						230					235		-		-	240
76 7	Tyr	Met	Ser	Lys	Ala	His	Gly	Val	Asp	Pro	Asn	Ile	Arq	Thr	Glv	Val
77	•			•	245		•		-	250			,		255	
79 I	Arq	Thr	Ile	Thr	Thr	Gly	Ser	Pro	Ile	Thr	Tyr	Ser	Thr	Tvr	Glv	Lvs
80	_			260		•			265		-			270		2
82 F	Phe	Leu	Ala	Asp	Gly	Gly	Cys	Ser	Gly	Gly	Ala	Tyr	qaA	Ile	Ile	Ile
83			275	-	•	•	•	280	4	- 4		- - -	285			
85 (Cys	Asp	Glu	Cys	His	Ser	Thr	qaA	Ala	Thr	Ser	Ile	Leu	Gly	Ile	Glv
86	•	290		•			295	-				300		_		
88 7	Thr	Val	Leu	Asp	Gln	Ala	Glu	Thr	Ala	Glv	Ala	Arq	Leu	Val	Val	Leu
89 3						310				4	315	J				320
91 <i>I</i>	Ala	Thr	Ala	Thr	Pro	Pro	Glv	Ser	Val	Thr	Val	Pro	His	Pro	Asn	
92					325		4			330					335	
94 (Glu	Glu	Val	Ala		Ser	Thr	Thr	Glv		Ile	Pro	Phe	Tvr	Glv	Lvs
95				340					345					350	1	-1-
97 <i>I</i>	Ala	Ile	Pro	Leu	Glu	Val	Ile	Lys	Gly	Gly	Arg	His	Leu	Ile	Phe	Cvs
98			355					360	•	-			365			-
100	His	Ser	Lys	Lys	Lys	Cys	Asr	Glu	Let	ı Ala	a Ala	Lys	Leu	ı Va]	Ala	Leu
101		370		•	•	•	375					380				
103	Gly	Ile	Asn	ı Ala	a Val	. Ala	. Tyr	Tyr	Arc	r Gly	/ Leu			. Sei	. Val	. Ile
104						390		4	_	, - 1	395					400
			Asn	Gly	Asp	Val	. Val	. Val	. Val	. Ala	Thr	Asr	Ala	Lei	ı Met	Thr
107				•	405					410		•			415	
109	Gly	Phe	Thr	Gly	/ Asp	Phe	Asp	Ser	. Val	. Ile	Asp	Cys	Asr	Thi	Cys	. Val
110	-			420			-		425		-	_		430	_	
112	Thr	Glr	Thr	. Val	. Asp	Phe	Ser	Leu	ı Asp	Pro	Thr	Phe	Thr	Ile	e Glu	1 Thr
113			435		_			440					445			
115	Thr	Thr	Leu	Pro	Glr	ı Asp	Ala	. Val	Ser	Arg	Thr	Glr	Arg	Arg	g Gly	/ Arg
116		450					455					460		-	_	_
118	Thr	Gly	Arg	Gly	/ Lys	Pro	Gly	r Ile	. Tyr	Arg	, Phe	Val	. Ala	Pro	Gly	/ Glu
119						470					475				_	480
121	Arg	Pro	Ser	Gly	/ Met	: Phe	. Asp	Ser	Ser	· Val	. Leu	Cys	Glu	Cys	Tyr	Asp
122					485		_			490		_		_	495	
124	Ala	Gly	y Cys	: Ala	Trp	Tyr	Gli	ı Lev	Thr	Pro	Ala	Glu	Thr	Thi	. Val	Arg
125				500		-			505					510		_
127	Leu	Arg	, Ala	Tyr	: Met	Asn	Thr	Pro	Gly	, Let	ı Pro	Val	. Cys	Glr	ı Asp	His
128			515	;				520)				525	;		
130	Leu	Gli	ı Phe	Trp	Glu	Gly	val	Phe	Thr	Gly	Leu	Thr	His	$_{ m I1}\epsilon$	a Asp	Ala
131		530		_		_	535			_		540			-	
133	His	Phe	e Leu	Ser	Glr	Thr	Lys	Glr	Ser	Gly	, Glu	Asr	Phe	Pro	туг	Leu
134						550	_			_	555				-	560
126																
T36	Val	Ala	а Туг	Glr	ı Ala	1 Thr	· Val	. Сув	: Ala	a Arg	, Ala	Glr	ı Ala	Pro	Pro	Pro

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139 Ser Trp Asp Gln Met Trp Lys Cys Leu Ile Arg Leu Lys Pro Thr Leu 140 580 585 142 His Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Ala Val Gln Asn Glu 595 600 143 605 145 Val Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Thr Cys Met Ser 615 148 Ala Asp Leu Glu Val Val Thr 149 625 630 153 <210> SEQ ID NO: 2 154 <211> LENGTH: 4 155 <212> TYPE: PRT 156 <213> ORGANISM: Artificial Sequence 158 <220> FEATURE: 159 <223> OTHER INFORMATION: Mutated Hepatitis C virus peptide 161 <400> SEQUENCE: 2 162 Ser Asp Gly Lys 163 1 167 <210> SEQ ID NO: 3 168 <211> LENGTH: 148 169 <212> TYPE: PRT 170 <213> ORGANISM: Hepatitis C virus 172 <400> SEQUENCE: 3 173 Gly Ser His Met Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala 176 Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 20 25 179 Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys 182 Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala 183 50 185 Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val 75 188 Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys 85 90 191 Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 100 105 194 Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly 115 120 197 Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 198 130 135 200 Ala Thr Ala Thr 201 145 205 <210> SEQ ID NO: 4 206 <211> LENGTH: 142 207 <212> TYPE: PRT 208 <213> ORGANISM: Hepatitis C virus 210 <400> SEQUENCE: 4 211 Gly Ser His Met Gly Ser Val Thr Val Pro His Pro Asn Ile Glu Glu 212

Input Set : A:\ID01152 Sequence Listing.txt
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214 Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys Ala Ile . 20 217 Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser 218 40 220 Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu Gly Ile 223 Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile Pro Thr 70 226 Asn Gly Asp Val Val Val Val Ala Thr Asp Ala Leu Met Thr Gly Phe 85 90 229 Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Ser Asp Gly Lys 100 105 232 Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr Gly Arg 115 120 235 Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu Arg 130 135 240 <210> SEO ID NO: 5 241 <211> LENGTH: 288 242 <212> TYPE: PRT 243 <213> ORGANISM: Hepatitis C virus 245 <400> SEQUENCE: 5 246 Gly Ser His Met Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala 247 1 10 249 Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 20 25 252 Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys 35 40 255 Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala 50 55 258 Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val 261 Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys 85 90 264 Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 105 267 Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly 115 120 270 Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 135 273 Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Val Pro His Pro Asn Ile 274 145 150 155 276 Glu Glu Val Ala Leu Ser Thr Thr Gly Glu Ile Pro Phe Tyr Gly Lys 170 279 Ala Ile Pro Leu Glu Val Ile Lys Gly Gly Arg His Leu Ile Phe Cys 280 180 185 282 His Ser Lys Lys Lys Cys Asp Glu Leu Ala Ala Lys Leu Val Ala Leu 195 200 205 285 Gly Ile Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ser Val Ile

Input Set : A:\ID01152 Sequence Listing.txt
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288 Pro Thr Asn Gly Asp Val Val Val Ala Thr Asp Ala Leu Met Thr 289 225 230 291 Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Thr Ser Asp 245 250 294 Gly Lys Pro Gln Asp Ala Val Ser Arg Thr Gln Arg Arg Gly Arg Thr 260 265 297 Gly Arg Gly Lys Pro Gly Ile Tyr Arg Phe Val Ala Pro Gly Glu Arg 298 275 280 305 <210> SEQ ID NO: 6 306 <211> LENGTH: 241 307 <212> TYPE: PRT 308 <213> ORGANISM: Hepatitis C virus 310 <400> SEQUENCE: 6 311 Gly Ser His Met Ser Pro Val Phe Thr Asp Asn Ser Ser Pro Pro Ala 312 1 5 10 314 Val Pro Gln Ser Phe Gln Val Ala His Leu His Ala Pro Thr Gly Ser 25 317 Gly Lys Ser Thr Lys Val Pro Ala Ala Tyr Ala Ala Gln Gly Tyr Lys 35 40 320 Val Leu Val Leu Asn Pro Ser Val Ala Ala Thr Leu Gly Phe Gly Ala 55 323 Tyr Met Ser Lys Ala His Gly Val Asp Pro Asn Ile Arg Thr Gly Val 70 75 $^{\circ}$ 326 Arg Thr Ile Thr Thr Gly Ser Pro Ile Thr Tyr Ser Thr Tyr Gly Lys 85 90 329 Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Ile Ile Ile 100 105 332 Cys Asp Glu Cys His Ser Thr Asp Ala Thr Ser Ile Leu Gly Ile Gly 115 120 335 Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu 130 135 338 Ala Thr Ala Thr Pro Pro Gly Ser Gly Met Phe Asp Ser Ser Val Leu 150 155 341 Cys Glu Cys Tyr Asp Ala Gly Cys Ala Trp Tyr Glu Leu Thr Pro Ala 165 170 344 Glu Thr Thr Val Arg Leu Arg Ala Tyr Met Asn Thr Pro Gly Leu Pro 180 185 190 347 Val Cys Gln Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu 348 195 200 350 Thr His Ile Asp Ala His Phe Leu Ser Gln Thr Lys Gln Ser Gly Glu 351 210 215 220 353 Asn Phe Pro Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Ala Arg Ala 354 225 235 240 356 Gln 361 <210> SEQ ID NO: 7 362 <211> LENGTH: 4 363 <212> TYPE: PRT 364 <213> ORGANISM: Artificial Sequence 366 <220> FEATURE:

VERIFICATION SUMMARY

DATE: 09/01/2004 TIME: 17:20:05

PATENT APPLICATION: US/09/825,423C

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